

(c) determining the nucleotide sequence of at least a portion of the DNA present in the amplified DNA sample.

In the abstract:

Please replace the abstract with the following version.

--The presently claimed invention concerns a method for biasing the amplification of the DNA molecules in a sample such that a nucleic acid molecule having a specific nucleotide at a selected position (e.g., a polymorphic site) is preferentially amplified relative to an otherwise identical nucleic acid molecule not having the specific nucleotide at the selected position. The method entails amplification of target nucleic acid molecules using a pair of primers one of which causes the incorporation into the amplification product of a sequence that interferes with amplification of nucleic acid molecules not having the specific nucleotide at the selected position. The method is useful for preferentially amplifying at least a portion of one allele of a gene relative to another, different allele of the gene in a sample containing both alleles of the gene.--

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